5

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:

Source:

"

Date Processed by STIC:

ENTERED



PCT

RAW SEQUENCE LISTING DATE: 03/31/2006
PATENT APPLICATION: US/10/520,158 TIME: 15:40:57

Input Set : F:\SEQUENCE LISTING.txt
Output Set: N:\CRF4\03312006\J520158.raw

```
3 <110> APPLICANT: Micronas Holding GmbH
              Klapproth, Holger
      6 <120> TITLE OF INVENTION: Composition and process for stabilizing of biomolecules
      8 <130> FILE REFERENCE: 1034
C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/520,158
C--> 11 <141> CURRENT FILING DATE: 2005-01-04
     13 <160> NUMBER OF SEQ ID NOS: 9
     15 <170> SOFTWARE: PatentIn Ver. 2.1
     17 <210> SEQ ID NO: 1
     18 <211> LENGTH: 11
     19 <212> TYPE: PRT
     20 <213> ORGANISM: Artificial sequence
     22 <220> FEATURE:
     23 <223> OTHER INFORMATION: Description of the artificial sequence: Motif of the LEA
class
     25 <220> FEATURE:
     26 <223> OTHER INFORMATION: Amino acid position 1: X= K or T
     28 <220> FEATURE:
     29 <223> OTHER INFORMATION: Amino acid position 2: X=A or G or K or M or T
     31 <220> FEATURE:
     32 <223> OTHER INFORMATION: Amino acid position 3: X=R or D or A or E or Q or K
     34 <220> FEATURE:
     35 <223> OTHER INFORMATION: Amino acid position 4: X= E or K or S
     37 <220> FEATURE:
     38 <223> OTHER INFORMATION: Amino acid position 5: X= T or F or Y or A
     40 <220> FEATURE:
     41 <223> OTHER INFORMATION: Amino acid position 6: X= K or R or T or A
     43 <220> FEATURE:
     44 <223> OTHER INFORMATION: Amino acid position 7: X= D or E or Q
     46 <220> FEATURE:
     47 <223> OTHER INFORMATION: Amino acid position 8: X= S or R or Y or K
     49 <220> FEATURE:
     50 <223> OTHER INFORMATION: Amino acid position 9: X= A or T
     52 <220> FEATURE:
     53 <223> OTHER INFORMATION: Amino acid position 10: X= G or A or R
     55 <400> SEQUENCE: 1
W--> 56 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Glu
     57
        1
                          5
                                             10
     60 <210> SEQ ID NO: 2
     61 <211> LENGTH: 11
     62 <212> TYPE: PRT
     63 <213> ORGANISM: Aphelenchus avenae
     65 <400> SEQUENCE: 2
     66 Lys Thr Ala Glu Phe Arg Asp Ser Ala Gly Glu
```

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```
67 1
                                        10
70 <210> SEO ID NO: 3
71 <211> LENGTH: 11
72 <212> TYPE: PRT
73 <213> ORGANISM: Aphelenchus avenae
75 <400> SEQUENCE: 3
76 Lys Gly Gln Glu Phe Lys Glu Arg Ala Gly Glu
77 1
80 <210> SEQ ID NO: 4
81 <211> LENGTH: 11
82 <212> TYPE: PRT
83 <213> ORGANISM: Aphelenchus avenae
85 <400> SEQUENCE: 4
86 Lys Ala Glu Glu Thr Lys Gln Arg Ala Gly Glu
87 1
90 <210> SEQ ID NO: 5
91 <211> LENGTH: 11
92 <212> TYPE: PRT
93 <213> ORGANISM: Aphelenchus avenae
95 <400> SEQUENCE: 5
96 Lys Met Asp Glu Thr Lys Gln Arg Ala Gly Glu
97 1
100 <210> SEQ ID NO: 6
101 <211> LENGTH: 11
102 <212> TYPE: PRT
103 <213> ORGANISM: Betula pendula
105 <400> SEQUENCE: 6
106 Lys Ala Arg Lys Thr Lys Asp Ser Ala Ala Glu
107
    1
                     5
110 <210> SEQ ID NO: 7
111 <211> LENGTH: 11
112 <212> TYPE: PRT
113 <213> ORGANISM: Betula pendula
115 <400> SEQUENCE: 7
116 Lys Ala Lys Glu Tyr Lys Asp Tyr Thr Ala Glu
117 1
                      5
120 <210> SEQ ID NO: 8
121 <211> LENGTH: 11
122 <212> TYPE: PRT
123 <213> ORGANISM: Betula pendula
125 <400> SEQUENCE: 8
126 Lys Ala Arg Glu Thr Thr Glu Lys Ala Arg Glu
127 1
130 <210> SEQ ID NO: 9
131 <211> LENGTH: 11
132 <212> TYPE: PRT
133 <213> ORGANISM: Betula pendula
135 <400> SEQUENCE: 9
136 Thr Lys Asp Ser Ala Ala Glu Lys Ala Arg Glu
```

RAW SEQUENCE LISTING

DATE: 03/31/2006

PATENT APPLICATION: US/10/520,158

TIME: 15:40:57

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137 1 5 10

RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/10/520,158

DATE: 03/31/2006 TIME: 15:40:58

Input Set : F:\SEQUENCE LISTING.txt
Output Set: N:\CRF4\03312006\J520158.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; Xaa Pos. 1/2/3/4/5/5/5/1/8/8/9/

VARIABLE LOCATION SUMMARY

PATENT APPLICATION: US/10/520,158

DATE: 03/31/2006 TIME: 15:40:58

Input Set : F:\SEQUENCE_LISTING.txt
Output Set: N:\CRF4\\\03312006\J520158.raw

Use of n's or Xaa's (NEW RULES):

Use of n's and/or Xaa's have been detected in the Sequence Listing.
Use of <220> to <223> is MANDATORY if n's or Xaa's are present.
in <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

Seq#:1; Xaa Pos. 1,2,3,4,5,6,7,8,9,10

VERIFICATION SUMMARYPATENT APPLICATION: US/10/520,158 DATE: 03/31/2006 TIME: 15:40:58

Input Set : F:\SEQUENCE LISTING.txt
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L:10 M:270 C: Current Application Number differs, Replaced Application Number L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:56 M:258 W: Mandatory Feature missing, <221> Tag not found for SEQ ID#:1 L:56 M:258 W: Mandatory Feature missing, <222> Tag not found for SEQ ID#:1 L:56 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0